SEQUENCE LISTING

<110> Rosanne M. Crooke
Mark J. Graham

<120> ANTISENSE MODULATION OF MICROSOMAL TRIGLYCERIDE TRANSFER PROTEIN EXPRESSION

<130> ISPH-0591

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Met Ile Leu Leu Ala Val Leu Phe Leu
1 5

tgc ttc att tcc tca tat tca gct tct gtt aaa ggt cac aca act ggt 161
Cys Phe Ile Ser Ser Tyr Ser Ala Ser Val Lys Gly His Thr Thr Gly
10 15 20 25

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ctc tca tta aat aat gac cgg ctg tac aag ctc acg tac tcc act gaa   209
Leu Ser Leu Asn Asn Asp Arg Leu Tyr Lys Leu Thr Tyr Ser Thr Glu
                 30                  35                  40

gtt ctt ctt gat cgg ggc aaa gga aaa ctg caa gac agc gtg ggc tac   257
Val Leu Leu Asp Arg Gly Lys Gly Lys Leu Gln Asp Ser Val Gly Tyr
             45                  50                  55

cgc att tcc tcc aac gtg gat gtg gcc tta cta tgg agg aat cct gat   305
Arg Ile Ser Ser Asn Val Asp Val Ala Leu Leu Trp Arg Asn Pro Asp
         60                  65                  70

ggt gat gat gac cag ttg atc caa ata acg atg aag gat gta aat gtt   353
Gly Asp Asp Asp Gln Leu Ile Gln Ile Thr Met Lys Asp Val Asn Val
     75                  80                  85

gaa aat gtg aat cag cag aga gga gag aag agc atc ttc aaa gga aaa   401
Glu Asn Val Asn Gln Gln Arg Gly Glu Lys Ser Ile Phe Lys Gly Lys
 90                  95                 100                 105

agc cca tct aaa ata atg gga aag gaa aac ttg gaa gct ctg caa aga   449
Ser Pro Ser Lys Ile Met Gly Lys Glu Asn Leu Glu Ala Leu Gln Arg
                110                 115                 120

cct acg ctc ctt cat cta atc cat gga aag gtc aaa gag ttc tac tca   497
Pro Thr Leu Leu His Leu Ile His Gly Lys Val Lys Glu Phe Tyr Ser
            125                 130                 135

tat caa aat gag gca gtg gcc ata gaa aat atc aag aga ggt ctg gct   545
Tyr Gln Asn Glu Ala Val Ala Ile Glu Asn Ile Lys Arg Gly Leu Ala
        140                 145                 150

agc cta ttt cag aca cag tta agc tct gga acc acc aat gag gta gat   593
Ser Leu Phe Gln Thr Gln Leu Ser Ser Gly Thr Thr Asn Glu Val Asp
    155                 160                 165

atc tct gga aat tgt aaa gtg acc tac cag gct cat caa gac aaa gtg   641
Ile Ser Gly Asn Cys Lys Val Thr Tyr Gln Ala His Gln Asp Lys Val
170                 175                 180                 185

atc aaa att aag gcc ttg gat tca tgc aaa ata gcg agg tct gga ttt   689
Ile Lys Ile Lys Ala Leu Asp Ser Cys Lys Ile Ala Arg Ser Gly Phe
                190                 195                 200

acg acc cca aat cag gtc ttg ggt gtc agt tca aaa gct aca tct gtc   737
Thr Thr Pro Asn Gln Val Leu Gly Val Ser Ser Lys Ala Thr Ser Val
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acc acc tat aag ata gaa gac agc ttt gtt ata gct gtg ctt gct gaa   785
Thr Thr Tyr Lys Ile Glu Asp Ser Phe Val Ile Ala Val Leu Ala Glu
        220                 225                 230

gaa aca cac aat ttt gga ctg aat ttc cta caa acc att aag ggg aaa   833
Glu Thr His Asn Phe Gly Leu Asn Phe Leu Gln Thr Ile Lys Gly Lys
    235                 240                 245

ata gta tcg aag cag aaa tta gag ctg aag aca acc gaa gca ggc cca   881
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Ile Val Ser Lys Gln Lys Leu Glu Leu Lys Thr Thr Glu Ala Gly Pro
250                 255                 260                 265

aga ttg atg tct gga aag cag gct gca gcc ata atc aaa gca gtt gat   929
Arg Leu Met Ser Gly Lys Gln Ala Ala Ala Ile Ile Lys Ala Val Asp
                270                 275                 280

tca aag tac acg gcc att ccc att gtg ggg cag gtc ttc cag agc cac   977
Ser Lys Tyr Thr Ala Ile Pro Ile Val Gly Gln Val Phe Gln Ser His
            285                 290                 295

tgt aaa gga tgt cct tct ctc tcg gag ctc tgg cgg tcc acc agg aaa   1025
Cys Lys Gly Cys Pro Ser Leu Ser Glu Leu Trp Arg Ser Thr Arg Lys
        300                 305                 310

tac ctg cag cct gac aac ctt tcc aag gct gag gct gtc aga aac ttc   1073
Tyr Leu Gln Pro Asp Asn Leu Ser Lys Ala Glu Ala Val Arg Asn Phe
    315                 320                 325

ctg gcc ttc att cag cac ctc agg act gcg aag aaa gaa gag atc ctt   1121
Leu Ala Phe Ile Gln His Leu Arg Thr Ala Lys Lys Glu Glu Ile Leu
330                 335                 340                 345

caa ata cta aag atg gaa aat aag gaa gta tta cct cag ctg gtg gat   1169
Gln Ile Leu Lys Met Glu Asn Lys Glu Val Leu Pro Gln Leu Val Asp
                350                 355                 360

gct gtc acc tct gct cag acc tca gac tca tta gaa gcc att ttg gac   1217
Ala Val Thr Ser Ala Gln Thr Ser Asp Ser Leu Glu Ala Ile Leu Asp
            365                 370                 375

ttt ttg gat ttc aaa agt gac agc agc att atc ctc cag gag agg ttt   1265
Phe Leu Asp Phe Lys Ser Asp Ser Ser Ile Ile Leu Gln Glu Arg Phe
        380                 385                 390

ctc tat gcc tgt gga ttt gct tct cat ccc aat gaa gaa ctc ctg aga   1313
Leu Tyr Ala Cys Gly Phe Ala Ser His Pro Asn Glu Glu Leu Leu Arg
    395                 400                 405

gcc ctc att agt aag ttc aaa ggt tct att ggt agc agt gac atc aga   1361
Ala Leu Ile Ser Lys Phe Lys Gly Ser Ile Gly Ser Ser Asp Ile Arg
410                 415                 420                 425

gaa act gtt atg atc atc act ggg aca ctt gtc aga aag ttg tgt cag   1409
Glu Thr Val Met Ile Ile Thr Gly Thr Leu Val Arg Lys Leu Cys Gln
                430                 435                 440

aat gaa ggc tgc aaa ctc aaa gca gta gtg gaa gct aag aag tta atc   1457
Asn Glu Gly Cys Lys Leu Lys Ala Val Val Glu Ala Lys Lys Leu Ile
            445                 450                 455

ctg gga gga ctt gaa aaa gca gag aaa aaa gag gac acc agg atg tat   1505
Leu Gly Gly Leu Glu Lys Ala Glu Lys Lys Glu Asp Thr Arg Met Tyr
        460                 465                 470

ctg ctg gct ttg aag aat gcc ctg ctt cca gaa ggc atc cca agt ctt   1553
Leu Leu Ala Leu Lys Asn Ala Leu Leu Pro Glu Gly Ile Pro Ser Leu
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    475                 480                 485

ctg aag tat gca gaa gca gga gaa ggg ccc atc agc cac ctg gct acc   1601
Leu Lys Tyr Ala Glu Ala Gly Glu Gly Pro Ile Ser His Leu Ala Thr
490                 495                 500                 505

act gct ctc cag aga tat gat ctc cct ttc ata act gat gag gtg aag   1649
Thr Ala Leu Gln Arg Tyr Asp Leu Pro Phe Ile Thr Asp Glu Val Lys
                510                 515                 520

aag acc tta aac aga ata tac cac caa aac cgt aaa gtt cat gaa aag   1697
Lys Thr Leu Asn Arg Ile Tyr His Gln Asn Arg Lys Val His Glu Lys
            525                 530                 535

act gtg cgc act gct gca gct gct atc att tta aat aac aat cca tcc   1745
Thr Val Arg Thr Ala Ala Ala Ala Ile Ile Leu Asn Asn Asn Pro Ser
        540                 545                 550

tac atg gac gtc aag aac atc ctg ctg tct att ggg gag ctt ccc caa   1793
Tyr Met Asp Val Lys Asn Ile Leu Leu Ser Ile Gly Glu Leu Pro Gln
    555                 560                 565

gaa atg aat aaa tac atg ctc gcc att gtt caa gac atc cta cgt ttg   1841
Glu Met Asn Lys Tyr Met Leu Ala Ile Val Gln Asp Ile Leu Arg Leu
570                 575                 580                 585

gaa atg cct gca agc aaa att gtc cgt cga gtt ctg aag gaa atg gtc   1889
Glu Met Pro Ala Ser Lys Ile Val Arg Arg Val Leu Lys Glu Met Val
                590                 595                 600

gct cac aat tat gac cgt ttc tcc agg agt gga tct tct tct gcc tac   1937
Ala His Asn Tyr Asp Arg Phe Ser Arg Ser Gly Ser Ser Ser Ala Tyr
            605                 610                 615

act ggc tac ata gaa cgt agt ccc cgt tcg gca tct act tac agc cta   1985
Thr Gly Tyr Ile Glu Arg Ser Pro Arg Ser Ala Ser Thr Tyr Ser Leu
        620                 625                 630

gac att ctc tac tcg ggt tct ggc att cta agg aga agt aac ctg aac   2033
Asp Ile Leu Tyr Ser Gly Ser Gly Ile Leu Arg Arg Ser Asn Leu Asn
    635                 640                 645

atc ttt cag tac att ggg aag gct ggt ctt cac ggt agc cag gtg gtt   2081
Ile Phe Gln Tyr Ile Gly Lys Ala Gly Leu His Gly Ser Gln Val Val
650                 655                 660                 665

att gaa gcc caa gga ctg gaa gcc tta atc gca gcc acc cct gac gag   2129
Ile Glu Ala Gln Gly Leu Glu Ala Leu Ile Ala Ala Thr Pro Asp Glu
                670                 675                 680

ggg gag gag aac ctt gac tcc tat gct ggt atg tca gcc atc ctc ttt   2177
Gly Glu Glu Asn Leu Asp Ser Tyr Ala Gly Met Ser Ala Ile Leu Phe
            685                 690                 695

gat gtt cag ctc aga cct gtc acc ttt ttc aac gga tac agt gat ttg   2225
Asp Val Gln Leu Arg Pro Val Thr Phe Phe Asn Gly Tyr Ser Asp Leu
        700                 705                 710
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atg tcc aaa atg ctg tca gca tct ggc gac cct atc agt gtg gtg aaa 2273
Met Ser Lys Met Leu Ser Ala Ser Gly Asp Pro Ile Ser Val Val Lys
715 720 725

gga ctt att ctg cta ata gat cat tct cag gaa ctt cag tta caa tct 2321
Gly Leu Ile Leu Leu Ile Asp His Ser Gln Glu Leu Gln Leu Gln Ser
730 735 740 745

gga cta aaa gcc aat ata gag gtc cag ggt ggt cta gct att gat att 2369
Gly Leu Lys Ala Asn Ile Glu Val Gln Gly Gly Leu Ala Ile Asp Ile
750 755 760

tca ggt gca atg gag ttt agc ttg tgg tat cgt gag tct aaa acc cga 2417
Ser Gly Ala Met Glu Phe Ser Leu Trp Tyr Arg Glu Ser Lys Thr Arg
765 770 775

gtg aaa aat agg gtg act gtg gta ata acc act gac atc aca gtg gac 2465
Val Lys Asn Arg Val Thr Val Val Ile Thr Thr Asp Ile Thr Val Asp
780 785 790

tcc tct ttt gtg aaa gct ggc ctg gaa acc agt aca gaa aca gaa gca 2513
Ser Ser Phe Val Lys Ala Gly Leu Glu Thr Ser Thr Glu Thr Glu Ala
795 800 805

ggc ttg gag ttt atc tcc aca gtg cag ttt tct cag tac cca ttc tta 2561
Gly Leu Glu Phe Ile Ser Thr Val Gln Phe Ser Gln Tyr Pro Phe Leu
810 815 820 825

gtt tgc atg cag atg gac aag gat gaa gct cca ttc agg caa ttt gag 2609
Val Cys Met Gln Met Asp Lys Asp Glu Ala Pro Phe Arg Gln Phe Glu
830 835 840

aaa aag tac gaa agg ctg tcc aca ggc aga ggt tat gtc tct cag aaa 2657
Lys Lys Tyr Glu Arg Leu Ser Thr Gly Arg Gly Tyr Val Ser Gln Lys
845 850 855

aga aaa gaa agc gta tta gca gga tgt gaa ttc ccg ctc cat caa gag 2705
Arg Lys Glu Ser Val Leu Ala Gly Cys Glu Phe Pro Leu His Gln Glu
860 865 870

aac tca gag atg tgc aaa gtg gtg ttt gcc cct cag ccg gat agt act 2753
Asn Ser Glu Met Cys Lys Val Val Phe Ala Pro Gln Pro Asp Ser Thr
875 880 885

tcc agc gga tgg ttt tga aactgacctg tgatatttta cttgaatttg 2801
Ser Ser Gly Trp Phe
890

tctccccgaa agggacacaa tgtggcatga ctaagtactt gctctctgag agcacagcgt 2861
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Cys Phe Phe Ser Ser Tyr Ser Ala Ser Val Lys Gly His Thr Thr Gly
10 15 20 25

ctc tca tta aat aat gag cgg cta tac aag ctc acg tac tcc act gaa 147
Leu Ser Leu Asn Asn Glu Arg Leu Tyr Lys Leu Thr Tyr Ser Thr Glu
30 35 40

gtg ttt ctt gat ggg ggc aaa gga aaa ccg caa gac agc gtg ggc tac 195
Val Phe Leu Asp Gly Gly Lys Gly Lys Pro Gln Asp Ser Val Gly Tyr
45 50 55

aaa atc tca tct gat gtg gac gtt gtg tta ctg tgg agg aat cct gat 243
Lys Ile Ser Ser Asp Val Asp Val Val Leu Leu Trp Arg Asn Pro Asp
60 65 70

ggt gat gat gat caa gtg atc caa gtc acg ata aca gct gtt aac gtt 291
Gly Asp Asp Asp Gln Val Ile Gln Val Thr Ile Thr Ala Val Asn Val
75 80 85

gaa aat gcg ggt caa cag aga ggc gag aag agc atc ttc cag ggc aaa 339
Glu Asn Ala Gly Gln Gln Arg Gly Glu Lys Ser Ile Phe Gln Gly Lys
90 95 100 105

agt aca cct aag atc ata ggg aag gac aac ctg gag gct ctg cag aga 387
Ser Thr Pro Lys Ile Ile Gly Lys Asp Asn Leu Glu Ala Leu Gln Arg
110 115 120

ccc atg ctt ctt cat ctg gtc cgg ggg aag gtc aag gag ttc tac tcc 435
Pro Met Leu Leu His Leu Val Arg Gly Lys Val Lys Glu Phe Tyr Ser
125 130 135

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tat gaa aac gag cca gtg ggc ata gaa aat ctc aag aga ggc ttg gct   483
Tyr Glu Asn Glu Pro Val Gly Ile Glu Asn Leu Lys Arg Gly Leu Ala
        140                 145                 150

agc tta ttc cag atg cag cta agc tct gga act acc aac gag gta gat   531
Ser Leu Phe Gln Met Gln Leu Ser Ser Gly Thr Thr Asn Glu Val Asp
    155                 160                 165

atc tct ggg gat tgt aaa gtg acc tac cag gcc caa caa gac aaa gtg   579
Ile Ser Gly Asp Cys Lys Val Thr Tyr Gln Ala Gln Gln Asp Lys Val
170                 175                 180                 185

gtc aaa att aag gct ctg gat aca tgc aaa att gag cgg tct gga ttt   627
Val Lys Ile Lys Ala Leu Asp Thr Cys Lys Ile Glu Arg Ser Gly Phe
                190                 195                 200

aca acg gca aac cag gtg ctg ggc gtc agt tca aaa gcc aca tct gtc   675
Thr Thr Ala Asn Gln Val Leu Gly Val Ser Ser Lys Ala Thr Ser Val
            205                 210                 215

act acc tac aag ata gag gac agc ttt gtc acc gct gtg ctt gca gaa   723
Thr Thr Tyr Lys Ile Glu Asp Ser Phe Val Thr Ala Val Leu Ala Glu
        220                 225                 230

gag acc agg gct ttt gcc ttg aac ttc caa caa acc ata gca gga aaa   771
Glu Thr Arg Ala Phe Ala Leu Asn Phe Gln Gln Thr Ile Ala Gly Lys
    235                 240                 245

ata gtg tca aag cag aaa ttg gag ctg aag aca act gaa gcc ggc cca   819
Ile Val Ser Lys Gln Lys Leu Glu Leu Lys Thr Thr Glu Ala Gly Pro
250                 255                 260                 265

agg atg atc ccc ggg aag caa gtg gca ggt gta att aaa gca gtt gat   867
Arg Met Ile Pro Gly Lys Gln Val Ala Gly Val Ile Lys Ala Val Asp
                270                 275                 280

tcc aaa tac aaa gcc att ccc att gtg gga cag gtc ctc gag cgt gtc   915
Ser Lys Tyr Lys Ala Ile Pro Ile Val Gly Gln Val Leu Glu Arg Val
            285                 290                 295

tgc aaa gga tgc cct tct ctg gcg gag cac tgg aag tcc atc aga aag   963
Cys Lys Gly Cys Pro Ser Leu Ala Glu His Trp Lys Ser Ile Arg Lys
        300                 305                 310

aac ctg gag cct gaa aac ctg tcc aag gcc gag gct gtc cag agc ttc   1011
Asn Leu Glu Pro Glu Asn Leu Ser Lys Ala Glu Ala Val Gln Ser Phe
    315                 320                 325

ctg gcc ttc atc cag cac ctc cgg act tcg agg aga gaa gag atc ctc   1059
Leu Ala Phe Ile Gln His Leu Arg Thr Ser Arg Arg Glu Glu Ile Leu
330                 335                 340                 345

cag att ctg aag gca gag aag aaa gaa gtg ctc cct cag ctg gtg gat   1107
Gln Ile Leu Lys Ala Glu Lys Lys Glu Val Leu Pro Gln Leu Val Asp
                350                 355                 360
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gcc gtc acc tct gct cag act cca gac tcg cta gaa gcc atc ctg gac   1155
Ala Val Thr Ser Ala Gln Thr Pro Asp Ser Leu Glu Ala Ile Leu Asp
            365                 370                 375

ttt ttg gat ttc aaa agt gac agc agt atc ata ctc cag gaa agg ttc   1203
Phe Leu Asp Phe Lys Ser Asp Ser Ser Ile Ile Leu Gln Glu Arg Phe
        380                 385                 390

ctc tat gcc tgt ggc ttt gcc acc cac cct gat gaa gaa ctc cta cga   1251
Leu Tyr Ala Cys Gly Phe Ala Thr His Pro Asp Glu Glu Leu Leu Arg
    395                 400                 405

gcc ctc ctt agt aag ttc aaa ggt tcc ttt gca agc aac gac atc aga   1299
Ala Leu Leu Ser Lys Phe Lys Gly Ser Phe Ala Ser Asn Asp Ile Arg
410                 415                 420                 425

gag tcg gtt atg atc atc att gga gcc cta gtc agg aag ctg tgt cag   1347
Glu Ser Val Met Ile Ile Ile Gly Ala Leu Val Arg Lys Leu Cys Gln
                430                 435                 440

aat gaa ggc tgc aag ctc aag gca gtg gtg gaa gct aag aag ctg atc   1395
Asn Glu Gly Cys Lys Leu Lys Ala Val Val Glu Ala Lys Lys Leu Ile
            445                 450                 455

ctg gga gga ctt gaa aaa cca gag aag aaa gaa gac acc aca atg tac   1443
Leu Gly Gly Leu Glu Lys Pro Glu Lys Lys Glu Asp Thr Thr Met Tyr
        460                 465                 470

ctg ctg gcc ctg aag aat gcc ttg ctt ccc gaa ggc atc ccg ctc ctt   1491
Leu Leu Ala Leu Lys Asn Ala Leu Leu Pro Glu Gly Ile Pro Leu Leu
    475                 480                 485

ctg aag tat gct gag gct gga gaa ggg ccc gtc agc cac ctg gcc acc   1539
Leu Lys Tyr Ala Glu Ala Gly Glu Gly Pro Val Ser His Leu Ala Thr
490                 495                 500                 505

act gtt ctc cag aga tac gat gtc tcc ttc atc aca gat gag gtg aag   1587
Thr Val Leu Gln Arg Tyr Asp Val Ser Phe Ile Thr Asp Glu Val Lys
                510                 515                 520

aag acc ttg aac agg ata tac cac cag aat cgt aag gtt cat gag aag   1635
Lys Thr Leu Asn Arg Ile Tyr His Gln Asn Arg Lys Val His Glu Lys
            525                 530                 535

acg gtg cgc aca act gcc gct gct gtc atc tta aag aac cca tcc tac   1683
Thr Val Arg Thr Thr Ala Ala Ala Val Ile Leu Lys Asn Pro Ser Tyr
        540                 545                 550

atg gat gtg aag aac atc ctg ctg tcc att ggg gaa ctc ccg aaa gag   1731
Met Asp Val Lys Asn Ile Leu Leu Ser Ile Gly Glu Leu Pro Lys Glu
    555                 560                 565

atg aac aaa tac atg ctc acc gtt gtg caa gac atc ctg cat ttt gaa   1779
Met Asn Lys Tyr Met Leu Thr Val Val Gln Asp Ile Leu His Phe Glu
570                 575                 580                 585

atg cct gca agc aaa atg atc cgt cga gtt ctc aag gag atg gct gtt   1827
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Met Pro Ala Ser Lys Met Ile Arg Arg Val Leu Lys Glu Met Ala Val
                590                 595                 600

cac aat tat gac cgt ttc tcc aag agt gga tcc tct tct gcc tat act    1875
His Asn Tyr Asp Arg Phe Ser Lys Ser Gly Ser Ser Ser Ala Tyr Thr
            605                 610                 615

ggc tac gta gaa cgt agc ccc cgt gca gcg tcc aca tac agc ctt gac    1923
Gly Tyr Val Glu Arg Ser Pro Arg Ala Ala Ser Thr Tyr Ser Leu Asp
        620                 625                 630

atc ctt tac tct ggc tct ggc att ctg agg aga agt aac ctg aac atc    1971
Ile Leu Tyr Ser Gly Ser Gly Ile Leu Arg Arg Ser Asn Leu Asn Ile
    635                 640                 645

ttc cag tac atc aaa gga aca gag ctt cat ggt agt cag gtg gtg att    2019
Phe Gln Tyr Ile Lys Gly Thr Glu Leu His Gly Ser Gln Val Val Ile
650                 655                 660                 665

gaa gcc caa ggg ctg gaa ggc tta att gca gcc act cct gat gaa gga    2067
Glu Ala Gln Gly Leu Glu Gly Leu Ile Ala Ala Thr Pro Asp Glu Gly
                670                 675                 680

gag gag aac ctt gac tct tat gct ggc atg tca gcc atc ctg ttt gat    2115
Glu Glu Asn Leu Asp Ser Tyr Ala Gly Met Ser Ala Ile Leu Phe Asp
            685                 690                 695

gtt cag ctt agg cct gtc aca ttt ttt aat gga tac agt gat ttg atg    2163
Val Gln Leu Arg Pro Val Thr Phe Phe Asn Gly Tyr Ser Asp Leu Met
        700                 705                 710

tcc aaa atg ctg tcg gca tcc ggc gac cct gtc agc gtg gtg aaa ggg    2211
Ser Lys Met Leu Ser Ala Ser Gly Asp Pro Val Ser Val Val Lys Gly
    715                 720                 725

ctt att ctg tta ata gac cat tct cag gat att cag ctg caa tct gga    2259
Leu Ile Leu Leu Ile Asp His Ser Gln Asp Ile Gln Leu Gln Ser Gly
730                 735                 740                 745

cta aag gcc aat atg gag atc cag ggt ggt cta gct att gat att tct    2307
Leu Lys Ala Asn Met Glu Ile Gln Gly Gly Leu Ala Ile Asp Ile Ser
                750                 755                 760

ggt tca atg gaa ttc agt ctg tgg tat cgc gag tct aaa acc cga gtg    2355
Gly Ser Met Glu Phe Ser Leu Trp Tyr Arg Glu Ser Lys Thr Arg Val
            765                 770                 775

aaa aat cgg gtg gct gtg gtg ata acc agc gac gtc aca gtg gat gcc    2403
Lys Asn Arg Val Ala Val Val Ile Thr Ser Asp Val Thr Val Asp Ala
        780                 785                 790

tct ttt gtg aaa gct ggt ctg gaa agc aga gcg gag aca gag gct ggg    2451
Ser Phe Val Lys Ala Gly Leu Glu Ser Arg Ala Glu Thr Glu Ala Gly
    795                 800                 805

ctg gag ttc atc tcc aca gtg cag ttc tca cag tac ccg ttc ttg gtc    2499
Leu Glu Phe Ile Ser Thr Val Gln Phe Ser Gln Tyr Pro Phe Leu Val
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810                 815                 820                 825

tgc atg cag atg gac aag gct gaa gcc cca ctc agg caa ttc gag aca   2547
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Lys Tyr Glu Arg Leu Ser Thr Gly Arg Gly Tyr Val Ser Arg Arg Arg
            845                 850                 855

aaa gag agc cta gtg gcc gga tgt gaa ctc ccc ctc cat caa cag aac   2643
Lys Glu Ser Leu Val Ala Gly Cys Glu Leu Pro Leu His Gln Gln Asn
        860                 865                 870

tct gag atg tgc aac gtg gta ttc cca cct cag cca gaa agc gat aac   2691
Ser Glu Met Cys Asn Val Val Phe Pro Pro Gln Pro Glu Ser Asp Asn
    875                 880                 885

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Ser Gly Gly Trp Phe
890

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